

AMENDMENTS TO THE CLAIMS:

Claims 1-24, 31-36, 37-43, 44-49, 51-56 and 58 are canceled without prejudice or disclaimer. Claims 25 and 30 are amended. The following is the status of the claims of the above-captioned application, as amended.

Claims 1-24 (Cancelled.)

Claim 25. (Currently amended.) A variant of a parent alpha-amylase, comprising ~~an alteration selected from the group of alterations corresponding to a substitution of a K at an amino acid position selected from the group consisting of 118, 320 and 458 R118K, R320K or R458K~~ (using SEQ ID NO:12 for numbering), wherein the variant has alpha-amylase activity, and wherein the variant has at least 80% homology to ~~an amino acid sequence selected from the group consisting of SEQ ID NO:2, SEQ ID NO:4, SEQ ID NO:6, SEQ ID NO:8, SEQ ID NO:10, SEQ ID NO:12 and SEQ ID NO:13.~~

Claim 26. (Previously presented.) The variant of claim 25, comprising an alteration corresponding to R118K.

Claim 27. (Previously presented.) The variant of claim 25, comprising an alteration corresponding to R320K.

Claim 28. (Previously presented.) The variant of claim 25, comprising an alteration corresponding to R458K.

Claim 29. (Previously presented.) The variant of claim 25, comprising an alteration corresponding to R118K, R320K and R458K.

Claim 30. (Currently amended.) The variant of claim 25, wherein the parent alpha-amylase is ~~selected from the group consisting of *B. licheniformis* alpha-amylase, *B. stearothermophilus* alpha-amylase, *B. amyloliquefaciens* alpha-amylase, *Bacillus* sp. NCIB 12289 alpha-amylase, *Bacillus* sp. NCIB 12512 alpha-amylase, *Bacillus* sp. NCIB 12513 alpha-amylase, *Bacillus* sp. DSM 9357 alpha-amylase, *Bacillus* sp. DSMZ no. 12649 alpha-amylase, *Bacillus* sp. KSM AP1378 alpha-amylase, *Bacillus* sp. KSM K36 alpha-amylase and *Bacillus* sp. KSM K38 alpha-~~

| amylase.

Claims 31-36 (Cancelled.)

Claim 36. (Previously presented.) The variant of claim 25, wherein the variant has at least 80% homology with an amino acid sequence of SEQ ID NO:12.

Claims 37-43 (Cancelled.)

Claim 43. (Previously presented.) The variant of claim 25, wherein the variant has at least 90% homology with an amino acid sequence of SEQ ID NO:12.

Claims 44-49 (Cancelled.)

Claim 50. (Previously presented.) The variant of claim 25, wherein the variant has at least 95% homology with an amino acid sequence of SEQ ID NO:12.

Claim 51-56. (Cancelled.)

Claim 57. (Previously presented.) The variant of claim 25, wherein the variant has at least 97% homology with an amino acid sequence of SEQ ID NO:12.

Claim 58. (Cancelled.)